REMARKS

Claims 1-12 currently are pending. Claims 11 and 12 have been withdrawn from consideration. Claims 1, 4 and 9 currently have been amended.

Claims 1-10 are rejected under 35 USC § 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

The examiner stated that claims 1 and 9 appear to be missing the delta symbol before "6-desaturase." Applicants have checked the pending claims and claims 1 and 9 are not missing the delta symbol before "6-desaturase." The examiner may not have the correct set of claims. Please refer the applicants' previously preliminary amendment.

Claims 4 and 9 currently have been amended to recite the proper Markush claim language.

Claims 1 and 9 currently have been amended to recite " $\Delta 6$ -desaturase activity" instead of "enzymatic activity."

The examiner stated that as written it is unclear whether the claims are drawn to derivatives of both SEQ ID NOs 1 and 2 or only 2. We think the claim clearly sets out that these are "derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequences shown in SEQ ID NO: 2..."

Claims 1-10 are rejected under 35 USC § 112, first paragraph, as failing to comply with the written description requirement. The examiner believes that the

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specification does not give sufficient description of the genus which has at least 50

percent homology to SEQ ID NO: 1 and 2.

To overcome this rejection, applicants amend restrict the claims to derivatives of

SEQ ID NO: 1 and/or SEQ ID NO: 2 having at least 85% homology. Applicants believe

that given the knowledge general available such as homology algorithms algorithm of

Karlin and Altschul (Proc. Natl. Acad. Sci. USA 87:2264-2268, 1990), one of ordinary

skill in the art should be able to derive 85% homologs.

Please charge any shortage in fees due in connection with the filing of this

paper, including Extension of Time fees to Deposit Account No. 11-0345. Please credit

any excess fees to such deposit account.

Respectfully submitted,

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COMPLETE LISTING OF ALL CLAIMS IN THE APPLICATION

- 1. (currently amended) A process of preparing unsaturated fatty acids, which comprises introducing, into an organism, at least one isolated nucleic acid sequence encoding a polypeptide having Δ6-desaturase activity, selected from the group consisting of:
 - a) a nucleic acid sequence having the sequence shown in SEQ ID NO: 1,
 - b) nucleic acid sequences which, as a result of the degeneracy of the genetic code, are derived from the sequence shown in SEQ ID NO: 1,
 - c) derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequences shown in SEQ ID NO: 2 and have at least <u>85</u> 50% homology at the amino acid level without substantially reducing the enzymatic action of the polypeptides,
 - and culturing this organism, where the cultured organism contains at least 1 mol% of unsaturated fatty acids based on the total fatty acid content in the organism.
- 2. (previously presented) The process as claimed in claim 1, wherein the nucleic acid sequence is derived from a plant or algae.
- (previously presented) The process as claimed in claim 1, wherein the nucleic acid sequence is derived from Physcomitrella patens.
- 4. (currently amended) The process as claimed in claim 1, wherein the organism is an organism selected from the group consisting of bacterium, fungus, ciliate, algae,

- cyanobacterium, animal or and plant.
- 5. (previously presented) The process as claimed in claim 1, wherein the organism is a plant or algae.
- 6. (previously presented) The process as claimed in claim 1, wherein the organism is an oil crop.
- 7. (previously presented) The process as claimed in claim 1, wherein the cultured organism contains at least 5% by weight of unsaturated fatty acids based on the total fatty acid content in the organism.
- 8. (previously presented) The process as claimed in claim 1, wherein the unsaturated fatty acids are isolated from the organism.
- 9. (currently amended) A transgenic organism selected from the group consisting of plants, fungi, ciliates, algae, bacteria, cyanobacteria or and animals comprising at least one isolated nucleic acid sequence encoding a polypeptide with Δ6-desaturase activity, selected from the group consisting of:
 - a) a nucleic acid sequence having the sequence shown in SEQ ID NO: 1,
 - b) nucleic acid sequences which, as a result of the degeneracy of the genetic code, are derived from the sequence shown in SEQ ID NO: 1,
 - c) derivatives of the nucleic acid sequence shown in SEQ ID NO: 1 which encode polypeptides with the amino acid sequences shown in SEQ ID NO: 2 and have at least 85 50% homology at the amino acid level without substantially reducing the enzymatic Δ6 desaturase action of the

polypeptides.

- 10. (previously presented) A transgenic organism as claimed in claim 9, wherein the organism is a plant or algae.
- 11. (withdrawn) An oil, lipid or fatty acid or a fraction thereof, prepared by the process as claimed in claim 1.
- 12. (withdrawn) The use of the oil, lipid or fatty acid composition as claimed in claim 11 or of a transgenic organism in feed, foodstuffs, cosmetics or pharmaceuticals.